

Fig. 1

TIAVHHAEG APLEPHSFWRCPVGEPYLSSDPKISLLIGPSLLSGS TIS V L LSIGEAIYA SS L TQG A IGK Y VLQL YISLNSDMFP TIAIHHADG SPLDPHDFWRCPVGEPLLSNNPNISLLPGPSLLSGS TIS V L LSIGDAIYA SS L TQG A IGK Y VLQL YISLNSDMYP RITT. GP. LINNPSFIPTA TVD V T LVINDLIYA TS L TRG Q IGK Y VLQL YISLNSDLVP AALINDNRY NGINQFYFSIAEGRNLTH. GP. LINNPSFIPTA TVD T T FSLIKTHWC TH V LNG Q HVS N FVSM IIEPTSAGFP NLLHHDAAY NGINKFILDYAHDFSI. GH. PLNNPSFIPTA SPN T I FSLIKTHWC TH V LNG Q HVS N YLSM ILVQTASGYP ILVNDLRF NGINKFILDYAHDFSI. GH. PLNMPSFIPTA SPN T I FSLIKTHWC TH V LNG R HSH H YLAL VLATATGRI BLAVDLRF GGIGKELIVDNASDVTSFYPSA. FQEHLNFIPAP TGS T I FDMSATHYC TH V LSG R HSH H YLAL VLATATGRI BSS0 B6S1	DLNPVVSHTYDIND VIAAGTRGYQL SMPTVD RT SSEGIEDLUVEDLIKGSTKSHRYRNSEVDLDHPFSALY SV N IATEGSLI LG DLKPVISHTYDIND VIAAGTRGYQL SLPTVN TT SSEGIEDLUVEDLIKGKTKSHRCKNEDITFDHPFSAMY SV S IKIENTLI LG DLNPRISHTFNIND LALLNTDVYQL STPKVD RS ASSGIEDLIVLDIVNHDGSISTTRFKNNNISFDQPYAALY SV P IYYKGKII LG EFRTLKTLYLSDGV VYAIPGGCWMY FVSTQP RD FSAAPPEQRIIIMYNNDTIVERIINPPGVLDVWATLN GT S VYLGWVL PI IFRTMKTIKLSDGI VAATRS KE ATTDLAELRIAFYYNNDTFIERVISPTGLEGNWATLN GT S VYLGWVL PA MFKTLKIQYLSDGI VSATPLGCDMI SKVTET EE NSAVPTLMAHGRLGFDGQYHEKDLDVTTLFEDWVANY GV G SFIDGRVW SV BISS BISS BISS BISS BISS BISS BISS B	LITPLQGDTKCRTQG QQVSQDTCNE LKITWLGGKQVVNVLIRVNDYLSERPKIRVTTIPITQNYL.A. LLKLGDRVXI T LTTPLQGDTKCVTNR ANVNQSVCND.LKITWLKKRQVVNVLIRINNXLSDRPKIVVETIPITQNYL.A. LLKLGKKIYI (T LEHPINENAICNTTG PGKTQRDCNQ/SHSPWFSDRRWVNSIIVVDKGLNSIPKLKVWTISMRQNYW~SS GELLLGNKIYI (T VIKGTSIWNNQANKYFIPQWVAAL SQNQATQVQN KSSYSSWFGNRMIQSGILACPLRQDLTNBCLVLPFSNDQVLMCAN (SLLMYGDSVYY) (Q LIKGTPSYNKQSSRYFIPKHPNIT AGKSSEQAAA/RSSYVIRYHSNRLLQSAVLICPLSDMHTARCNLVWFNNQTLMCAN (LXVIDNNLXY Q LIKGTPSYNKQSSRYFIPKHPNIT AGKSSEQAAA/RSSYVIRYHSNRLLQSAVLICPLSDMHTARCNLVWFNNQTLMCAN (LXVIDNNLXY Q CVLRNSPSDTVQEGKYVIYKRYNDT PDEQDYQIRM~KSSYKPGRFGGKRIQQAILSIKVSTSLGEDPVLTVPPNTVTLMCAN (LTTVGTSHFL) (Q B332	SSGWHSQLQIGVLDVSHPLTINWTPHEALS NKE NWYNR RE IS TAYP SPDAANVATUT YANTS V TIM STSWHSKLQLGILDITOYSDIRIKWTWHNVLS NNE PWGHS DG IT TAYP NPTGSIVSSVI DSQKT V VIT SNSWHSKLQLGILDITOYSDIRIKWTWHNVLS NNE PWGHS DG IT TAYP NPTGSIVSSVI DSQKT V VIT SNSWWPMTMLYKVTITFINGQPSALSAQNVPTQQVP TGD SATNR GF LT A AWL TNPSSTSTFGSEATFTGSY NTATQ I TWY SSWWSASLFYRINTDFSKGIPPIIEAQWVPSYQVP VMP NATSF AN IT A VWP NDPEPTSQNALNPNYRFAGAF RNESN T TFY STSWWPYELLYEISFTFTNSGQSSVNMSWIPIYSFT SGN SGENV TA VS I PWP TPYSHQSGINRNFYFTGAL NSSTT V TLY STSWWPYELLYPMTVSNKTATLHSPTTFNAFT SIP QASAR NS VT T PYP IFYRNHTLRGVFGTM DSEQA L ASA	YSNTTNINMERIKDVQLEVA TTISS ITHFGKG., FH I INQKSLNTLQPMLFKTSIPKLCKAES YSNTSEINMERKNVQLEAA TTTS. ITHFGKG., FH V INQTSLNTLQPMLFKTSIPKICKITS YSTATERVNELAIRNKTLSAG TTTS. ITHFKKG., FH V INHKSLDTFQPMLFKTEIPKSCS IANNTQLISTSQQEGSSGQEAA GHTT. FRDTGSVMV IY I LSSSLLGQFQIVPFIRQVTLS TASASALLNTTGFRNTUHKAA TSST. FKNTGTQKI LI I MGSSLLGGFQIIPFLRELIP VSALNNLKVLAPYGNQGLFAS TTTT. FQDTGDASV VY M LASNIVGEFQILPVLTRLTIT VSALNNLKVLAPYGNQGLFAS TTTT. FVVKTNKT LS A ISNTLFGEFRIVPLLVEILKNDGVREARSG 555
Sendai PIV1 PIV3 SV5 PIV2 Mumps NDV (12	Sendar PIV1 PIV3 SV5 PIV2 Mumps	sendai PIV1 PIV1 VIV3 SV5 PIV2 Mumps NDV (31	Sendai PIV1 PIV3 SV5 PIV2 Mumps	Sendar PIV1 PIV3 SV5 PIV2 Mumps NDV (50

Figure 2

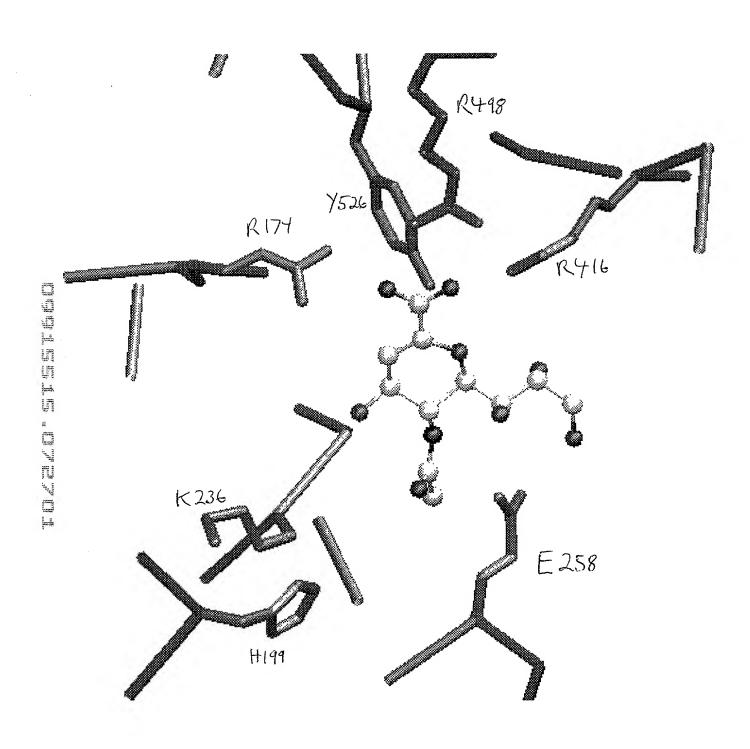


Fig. 3

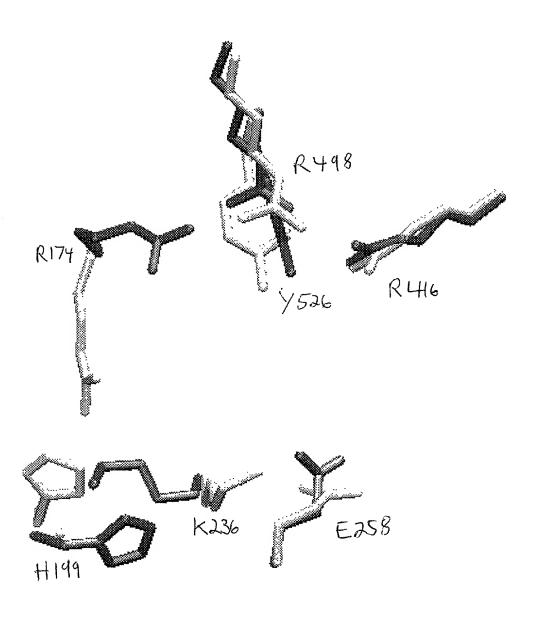


Fig. 4

	Native 1	Native 2	Native 3	Native 4	NANA	DANA
Cell (Å)	73.3	72.3	71.7	72.0	71.6	137.5
	78.0	77.9	77.9	83.8	77.6	137.5
	202.6	199.2	198.2	201.5	197.2	116.6
Temp(K),	293, 7	100, 4.6	100, 4.6	293, 6	100K, 4.6	100K,6.5
Resolution	າ 3.0	2.0	2.5	3.0	2.5	2.8
# obs	172104	623166	420703	277932	210555	498619
# unique	20022	68217	38168	22207	29671	38673
Complete ness(%)	83	86	97	88	76	94
R _{merg} (%)	9.3	4.9	3.1	9.3	4.2	5.2
R-factor			0.222		.223	0.209
R_{free}			0.277		.291	0.235
# protein			6914		6914	6896
atoms		·				
# CHO, Ca,			89		116	111
υa,			_			

Fig. 5

ligands

# waters	211	207	239
 A, B Å²	25,36	32,44	44,44
 ligand Å²	•	38,57	48,41

Table 1 Crystallographic data and refinement statistics.

Datasets Native2, Native3, NANA and DANA from frozen crystals were collected on beamlines X11 and BW7A at DESY, Hamburg. All other datasets were collected on in-house rotating anode and image plate or multiwire detector systems.

Remerge = $\sum_{hkl} \sum_{i} |\sum_{hkl} i'_{hkl} - \langle i_{hkl} \rangle| / \sum_{hkl} \sum_{i} \langle i'_{hkl} \rangle$ where the sum i is over all separate reasurements of the unique reflections hkl.

$$R$$
-factor = Σ_{hkl} $||F_{obs}| - |F_{calc}|| / \Sigma_{hkl} ||F_{obs}||$

R_{free}, as R-factor but summed over a 10% test set of reflections.